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3D1 HEAVY CHAIN VARIABLE REGION SEQUENCE

GCA A V GIC ATG M C AAG K CIT AAG K CAG Q မရှိ မရှိ GCC A FIL GGT TOTO AGT S TGG CTA L S AGA R CAG Q AAC N AAG K $_{\rm C}^{\rm TGT}$ 9 9 9 TGG W 99 GAG E CAG Q TIG ATC I GGT ACA T AAG K TAC Y TCT TGG W CAA Q GCC A ACA T 999 999 ATC I ATT STOT GAG E GGA G ACA T GGA G TTC Č T TTC GAT D V V ACC TCA T S GAG E ATG M ACT T TTT 150 T GAT TAT GCT ATA D Y A I 30 CTG L 270 ACT T 210 ATT I OTG OTG 330 TCT S GTC ACC GTA V AAT N $^{\rm GTT}_{\rm V}$ GCC A GTG V ATC I ATT GAC D ACA T AGG R TAC AAA K CCT ACA T TAT Y 999 9 S TAT Y CAG TCC TAC Y GCT A TGG W GAT D GAA E TGT S ACA T TCA S GTG V GCA A AGC S AAT GGT GGT TCA AAG K ACA T ACA T AGA R GTG V GIG V GCG A CAG Q GCC A AAC N AAG K CAC H \Pr_{Y}^{TAT} GCC A AGT S ATT S TGG 300 ATG 240 AAC N

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NAMST-JARIO BA CLASS SUBCLASS **APPROVED** O.G. FIG.



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3<u>D</u>1 LIGHT CHAIN VARIABLE REGION SEQUENCE

GAC □ ATG M ATG M TGG W AGC S TAC Y GAT D AGC S ATT CAG Q 999 999 GIG V TCA S AGT S TGC CAG AAA CCA Q K P QTC V AAA K_ CIG CAG Q GIG V TCC AGT CAG GCC CAG A Q CCT TCA S CAG Q GAT D CAG Q GCT A GTT CGC R 9 999 GAA E TCT AGT S CTT CAG Q CCA P GAC D TTC STCT ACA T CIG TG ATA I S DOL 30 TTG 210 CCT 330 GCA A 270 GGC G 150 CTC 90 TCC S GTT TAT V Y AAA K CTG AGT S TOTO AAC N GGA G T CTG AGT S GCT A T CIG CTG AGA R TAC Y GTG V CTA L STOT ATC I 999 999 TCA S DOL JOI ACC T TGG W TAC TGG GCA TCC ACT .
Y W A S T
ACA GAT TTC ACT CTC . CGA R GTA V GCA A ACG CAA TCT TAT AAT
T O S Y N GAG AAC E N GGA G STOT GAG E e GGT TAC AAG K ACC T TIG GTC V. CI 120 ACT T ი მემ 00 300 ACC T 180 GCT A

TTC

GGA G

999 999

99 99

ACC T

AAG K

CTG GAA L E

ATA AAA I K **1(B)**

NAMŻTJARO 趴 CLASS SUBCLASS APPROVED O.G. FIG.



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Hu3D1 HEAVY CHAIN VARIABLE REGION SEQUENCE

GAA E	CAG Q	GGA G	00 TGC	GTC V	ATG M
CTT	AAG K	CAG Q	AAA K	CAG Q	GGT G
AGT S	TTT	9 9 9	GCT A	CIG	TGG W
S	AAG K	TCTC	S	$\operatorname*{GTG}_{V}$	AAC N
TTG	9 299	GAG E	9 9 9	CAG Q	TGT
AGA R	AAG K	TGG W	TAC Y	STOT	ATC I
STOT	GCC A	ATT I	ACA T	6 6 6	ATC I
GAG E	ACA T	GGA G	TTC	GCT A	TTC
GAT D	ATG M	GTT V_	ACT	GAG E	TTT
330 ACG T	ACT T	210 ATT	150 GAT D	90 GTG V	30 CTG L
GCC A	GTA V	AAT N	TAT Y	AAG K	GTT V
$^{\tt GTT}_{\tt V}$	GAC D	ATT	GCT A	AAG K	ACC T
TAT	AAG K	TAC Y	ATA	CCI	ACA T
TAC Y	TCG	TAT Y	CAG Q	G G	GCT A
TGT	ACG T	GAT D	TGG W	AGC S	ACA T
GCA A	AGC S	AAT N	GTG V	TCA S	G G T
AGA R	ACA T	ACA T	AGA R	$_{\rm V}^{\rm GTG}$	OTG V
GCG A	GCC A	AAC N	CAG Q	AAG K	CAC H
GCC A	$_{Y}^{\mathtt{TAT}}$	TAC Y	GCT A	GTG V	S
360 TGG W	ATG M	240 AAC N	180 CCT P	120 TCC S	CAG

FIG. 2(A)

TAT ATG GAC TAC TGG GGT CAA GGT ACC CTT GTC ACC GTC TCC TCA
Y M D Y W G Q G T L V T V S S

NAMETTARIO 묎 APPROVED O.G. FIG.

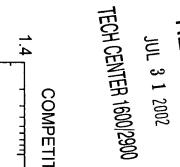


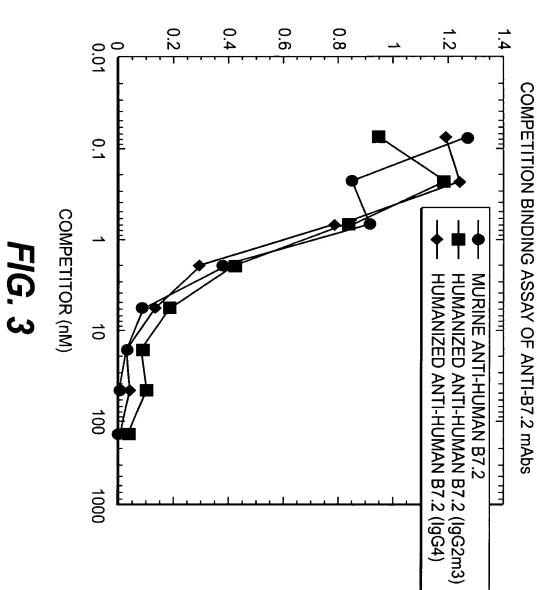
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CHAIN	
Hu3D1 LIGHT CHAIN VARIABLE	
REGION S	
SEQUENCE	

TAC P	ATC I	GAA E	TGG W	ATT	GAC <u>D</u>	ATG M
ACG T	AGC S	STOT	TAC Y	AGC S	ATT	·GAT D
TTC	AGT S	9 999	CAG Q	TGC AAA '	$_{\rm V}^{\rm GTG}$	TCA S
GGA G	CTG	GTC V	CAG Q	AAA K	CTG	CAG Q
CAG Q	CAG Q	CCT	AAA K	TCC AGT	ACA T	GCC A
9 9	GCT A	GAT D	CCA P	AGT S	CAG Q	CAG Q
ACC T	GAA E	CGC TTC	9 9 9	CAG AGT	TCT	$_{\rm V}^{\rm GTT}$
AAG K	GAC D	TTC	CAG Q	AGT	CCA P	LIL
T ∨ GIIG	V	AGT S	CCT	CIG	GAT D	ATA I
990 GAA E	330 GCA A	270 GGC G	210 CCT P	150 CTC	90 TCC S	30 TTG L
ATA I	GTT V	AGT S	AAA K	N AAC A	CIG	CTG (
	TAT Y	GGA G	T CIG	S GI	GCT A	CTG
	TAC Y	TCT	CTG	AGA R	GTA V	CTA L
	TGC	9 999	CTG ATC	AGA ACC R	AGC S	TGG W
	ACG	ACA T	TAC Y	CGA R	TTA L	$_{\rm V}^{\rm GTA}$
	CAA O	GAT D	TGG W	GAG E	GGA G	STOT
	TCT	TTC	GCA A	AAC N	GAG E	9 9 9
	TAT	ACT T	TCC	TAC Y	AGG R	ACC T
	AAT N	GAT TTC ACT CTC ACC D F T L T	TGG GCA TCC ACT AGG	GAG AAC TAC TTG GCT E N Y L A	GCC A	TGT
	CAA TCT TAT AAT CTT	300 ACC T	240 AGG R	180 GCT A	120 A GAG AGG GCC ACT E R A T	GGC ACC TGT GGG G T C G







BOUND/FREE

PPPROVED O.G. FIG.

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DIRECT BINDING ASSAY OF ANTI-B7.2 mAbs

ANTIBODY BOUND (cpm)

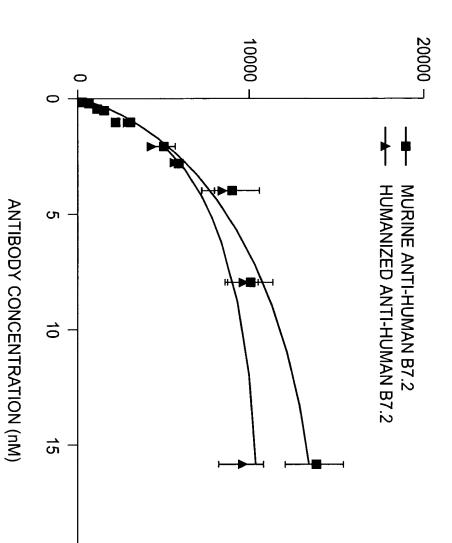


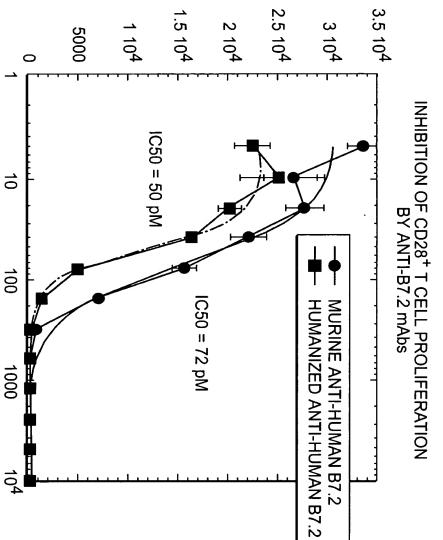
FIG. 4

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		NAMSTHARO
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IC.	O.G. F	DBVOR99A



$[^3$ H]-THYMIDINE INCORPORATION (CPM)



mAb CONCENTRATION [pM]

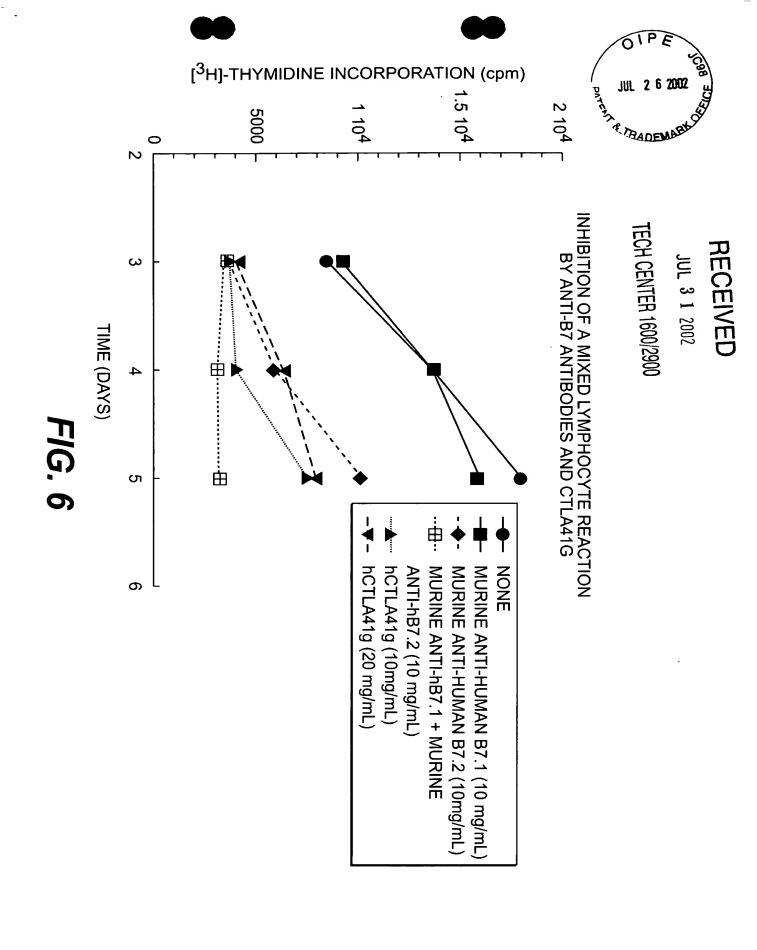
FIG. 5

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BY CLASS SUBCLASS

DRAFTSMAN



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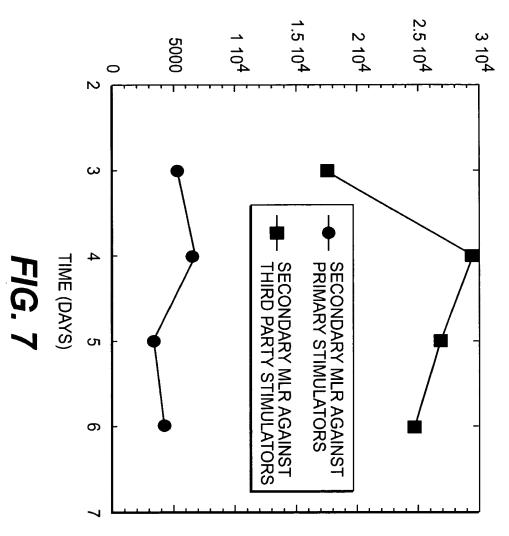
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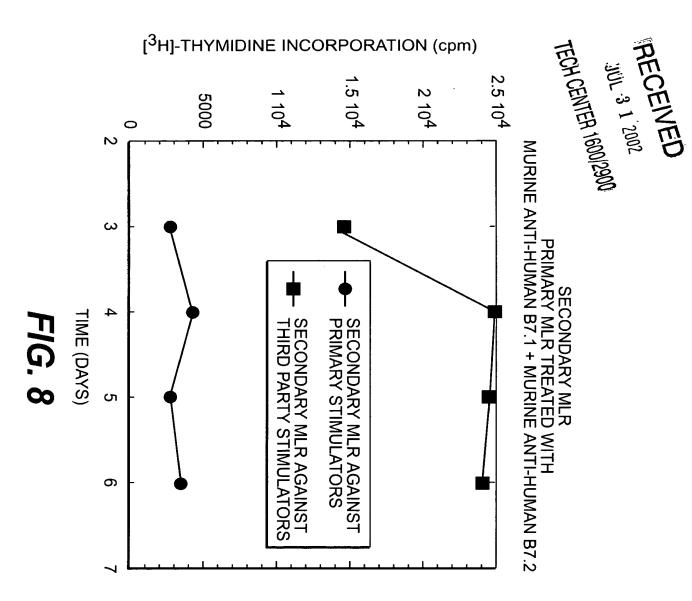
SECONDARY MLR
PRIMARY MLR TREATED WITH MURINE ANTI-HUMAN B7.2

[³H]-THYMIDINE INCORPORATION (cpm)

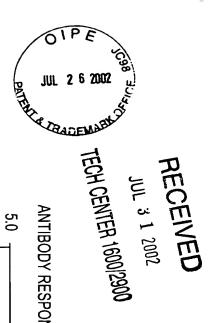


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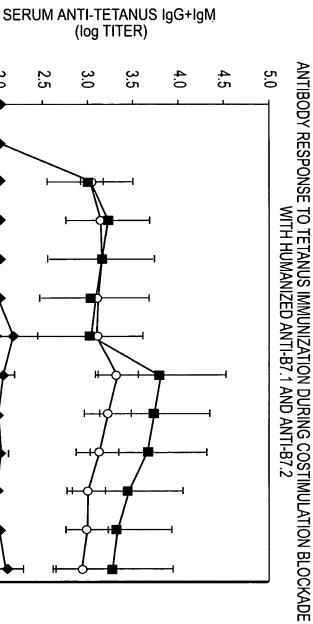


FIG. 9

GROUP D (10 mg/kg ANTI-B7 mAbs DAYS 0 AND 42)

TETANUS:

<u>.</u>5

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28

35

42

49

56

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70

77

22

2.0

■■ GROUP A (NO ANTI-B7 mAbs)

TIME (DAYS)

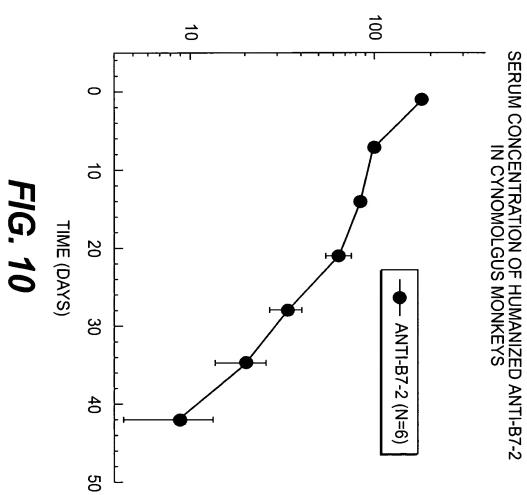
GROUP B (10 mg/kg ANTI-B7 mAbs DAY 0)
GROUP C (10 mg/kg ANTI-B7 mAbs DAY 42)

РРРПО О.G. FIG.

ANTI-B7-2, SERUM CONCENTRATION (ug/mL)



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